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APPLICATION NO. FILING DATE FIRST NAMED INVENTOR

TO THE PROPERTY OF THE PROPE

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DATE MAILED:

ART UNIT

64768 PM

PAPER NUMBER

Please find below and/or attached an Office communication concerning this application or proceeding.

Commissioner of Patents and Trademarks

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SERIAL NUMBER	FILING DATE	FIRST NAMED APPLICANT		ATTORNEY DOCKET NO.
09 004395 p	01:08/98	GILMORE ET AL		97 429
				EXAMINER
		11M	NNIFIELD	
			ART UNIT	PAPER NUMBER
		164	45	8
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Commissioner of Patents

Please find below a communication from the EXAMINER in charge of this application.

1. The reply filed January 15, 1999 and March 18, 1999 is not fully responsive to the communication mailed December 7, 1998 for the reasons set forth on the attached Notice To Comply With The Sequence Rules or CRF Diskette Problem Report.

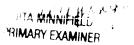
Since the above-mentioned reply appears to be *bona fide*, applicant is given a TIME PERIOD of **ONE (1) MONTH** or **THIRTY (30) DAYS**, from the mailing date of this notice, whichever is longer, within which to supply the omission or correction in order to avoid abandonment. EXTENSIONS OF THIS TIME LIMIT MAY BE GRANTED UNDER 37 CFR 1.136(a).

2. Any inquiry concerning this communication or earlier communications from the examiner should be directed to N. M. Minnifield whose telephone number is (703) 305-3394. The examiner can normally be reached on Monday-Thursday from 7:00 AM-4:30 PM. The examiner can also be reached on alternate Fridays

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor. Anthony Caputa, Ph.D., can be reached on (703) 308-3995. The fax phone number for this Group is (703) 308-4556.

Any inquiry of a general nature or relating to the status of this application should be directed to the Group receptionist whose telephone number is (703) 308-0196.

March Cran



Raw Sequence Listing Error Summary

٠	ERROR DETECTED	SUGGESTED	CORRECTION	SERIAL NUMBER: CY/OC4, 395		
ATTA	: NEW RULES CASES:	PLEASE DISREGA	ARD ENGLISH "ALPHA" H	EADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
	Wrapped Nucleics			pped" down to the next line.		
·	,,			word processor after creating it.		
			our right margin to .3, as this			
2	Wrapped Aminos	The amino acid r	number/text at the end of ea	ch line "wrapped" down to the next line.		
This may occur if your file was retrieved in a word processor after cre						
			our right margin to .3, as this			
3	Incorrect Line Length	The rules require	that a line not exceed 72 c	haracters in length. This includes spaces.		
	•	All text must be		·-		
4	Misaligned Amino Acid	. The numbering u	nder each 5th amino acid is	misaligned. This may be caused by the use of tabs		
· ——	Numbering	between the numbering. It is recommended to delete any tabs and uses spacing between the numbers.				
5	Non-ASCII	This file was not s	saved in ASCII (DOS) text,	as required by the Sequence Rules.		
		Please ensure your subsequent submission is saved in ASCII text so that it can be processed.				
6	Variable Length	Sequence(s)	_contain n's or Xaa's which	represented more than one residue.		
		As per the rules, e	each n or Xaa can only repr	esent a single residue.		
		Please present the maximum number of each residue having variable length and				
		indicate in the (ix)) features section that some	may be missing.		
7	Wrong Designation	Sequence(s)	contain amino acid or nucl	eic acid designators which are not standard		
		representations as	per the Sequence Rules (I	Please refer to paragraph 1.822)		
	Skipped Sequences			se use the following format for each skipped sequence:		
	(OLD RULES)	(2) INFORMATION	N FOR SEQ ID NO:X:			
				insert any headings under "SEQUENCE CHARACTERISTICS")		
		• •	ESCRIPTION:SEQ ID NO:	X:		
		This sequence is	Intentionally skipped			
		Please also adjust	the "(iii) NUMBER OF SEC	UENCES:" response to include the skipped sequence(s).		
9 :	Skipped Sequences	Sequence(s)	missing. If Intentional, plea	se use the following format for each skipped sequence.		
((NEW RULES)	<210> sequence	ld number			
		<400> sequence	ld number			
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10 l	Use of N's or Xara's	Use of N's and/or X	(aa's have been detected in	the Sequence Listing.		
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12 0 0	Jse of <220>Feature	Sequence(s)	are missing the <220>Featu	ore and associated headings.		
		e i same per esta esta esta esta esta esta esta esta	empte: 5%			
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3 W	Vrong Format	File submitted was t	n the alphabetical heading	format of the Old Sequence Rules.This is invalid since the		
		Requirements for P	atent Applications Containi	ng Nucleotide Sequence and/or Amino Acid Disclosures*		
			tice, Vol. 63, No. 104, June			
			ns filed on or after July 1, 1			

AKS Biotechnology Systems Branch: 7/50/19

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